

CRF Errors Corrected by the STIC Systems Branch

0400 124.01 01/PE
 CRF Processing Date: 1/10/2001
 Edited by: [Signature]
 Verified by: [Signature] (STIC staff)

Serial Number: 09/746,375

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically:
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form. 3/1/95

OIPE

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/746,375

DATE: 01/10/2001
 TIME: 11:25:17

Input Set : A:\99-106.SEQ.txt
 Output Set: N:\CRF3\01102001\I746375.raw

**Does Not Comply
 Corrected Diskette Needed**

4 <110> APPLICANT: Presnell, Scott, R.
 5 Kindsvogel, Wayne
 7 <120> TITLE OF INVENTION: NOVEL CYTOKINE ZCYTO18
 9 <130> FILE REFERENCE: 99-106
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/746,375
 C--> 11 <141> CURRENT FILING DATE: 2000-12-22
 11 <150> PRIOR APPLICATION NUMBER: US 60/172,105
 12 <151> PRIOR FILING DATE: 1999-12-23
 14 <150> PRIOR APPLICATION NUMBER: US 60/***,***
 15 <151> PRIOR FILING DATE: 2000-12-01
 17 <160> NUMBER OF SEQ ID NOS: 44
 19 <170> SOFTWARE: FastSEQ for Windows Version 3.0

ERRORED SEQUENCES

1031 <210> SEQ ID NO: 44
 1032 <211> LENGTH: 18
 1033 <212> TYPE: DNA
 1034 <213> ORGANISM: Artificial Sequence
 1036 <220> FEATURE:
 1037 <223> OTHER INFORMATION: Oligonucleotide primer ZC976
 1039 <400> SEQUENCE: 44
 1040 cgttgtaaaa cgacggcc
 E--> 1046 25

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VERIFICATION SUMMARY DATE: 01/10/2001
PATENT APPLICATION: US/09/746,375 TIME: 11:25:19

Input Set : A:\99-106.SEQ.txt
Output Set: N:\CRF3\01102001\I746375.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:162 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:163 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:164 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:165 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:166 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:167 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:168 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:169 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:170 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:1046 M:254 E: No. of Bases conflict, LENGTH:Input:25 Counted:18 SEQ:44